

Table S1 Ninety DEGs extracted from TCGA database had the significant value on prognosis of OSCC patient

Order number	Gene symbol
1	GZMK
2	ADAMDEC1
3	PLA2G2D
4	IGHV3-73
5	IGLV3-21
6	OR2I1P
7	MS4A1
8	CXCR3
9	IGHV3-33
10	FASLG
11	UBD
12	IGHV3-30
13	IGHGP
14	TRBV19
15	IGHV3-15
16	GZMA
17	NKG7
18	IGLL5
19	FCRL3
20	IGHM
21	TRBV5-1
22	IGKV5-2
23	IGHG2
24	IDO1
25	GZMM
26	SIRPG
27	IGHV1-69
28	SH2D1A
29	CCR5
30	TRAV21
31	CD40LG
32	TRBV4-2
33	IGHG3
34	TRBC2

Table S1 (continued)**Table S1** (continued)

Order number	Gene symbol
35	IGHG1
36	IGHV4-31
37	LINC00861
38	IKZF3
39	TRBV28
40	IGLV6-57
41	LY9
42	MYBPC1
43	CD2
44	IGLV7-43
45	CD3E
46	TRAV17
47	IGHG4
48	TIGIT
49	AC015911.7
50	TIFAB
51	FCRL5
52	GPR25
53	SAMD3
54	CXCR6
55	WDFY4
56	CD79A
57	IL12RB1
58	VCAM1
59	SIT1
60	CD5
61	ABCD2
62	BTLA
63	RUFY4
64	LINC00426
65	SPOCK2
66	CCR7
67	TRAC
68	ADH1B

Table S1 (continued)

Table S1 (*continued*)

Order number	Gene symbol
69	<i>IL9R</i>
70	<i>IGKV3D-11</i>
71	<i>IGHJ3</i>
72	<i>CD7</i>
73	<i>IRF8</i>
74	<i>GPR171</i>
75	<i>CLECL1</i>
76	<i>TRAV8-6</i>
77	<i>KLRD1</i>
78	<i>GRAP2</i>
79	<i>ZAP70</i>
80	<i>CHRM3-AS2</i>
81	<i>CLEC9A</i>
82	<i>CTLA4</i>
83	<i>CD96</i>
84	<i>NCR1</i>
85	<i>CD28</i>
86	<i>FAM30A</i>
87	<i>TBC1D10C</i>
88	<i>TNFRSF17</i>
89	<i>AC105460.2</i>
90	<i>AC105460.1</i>

DEGs, differentially expressed genes; TCGA, The Cancer Genome Atlas; OSCC, oral squamous cell carcinoma.